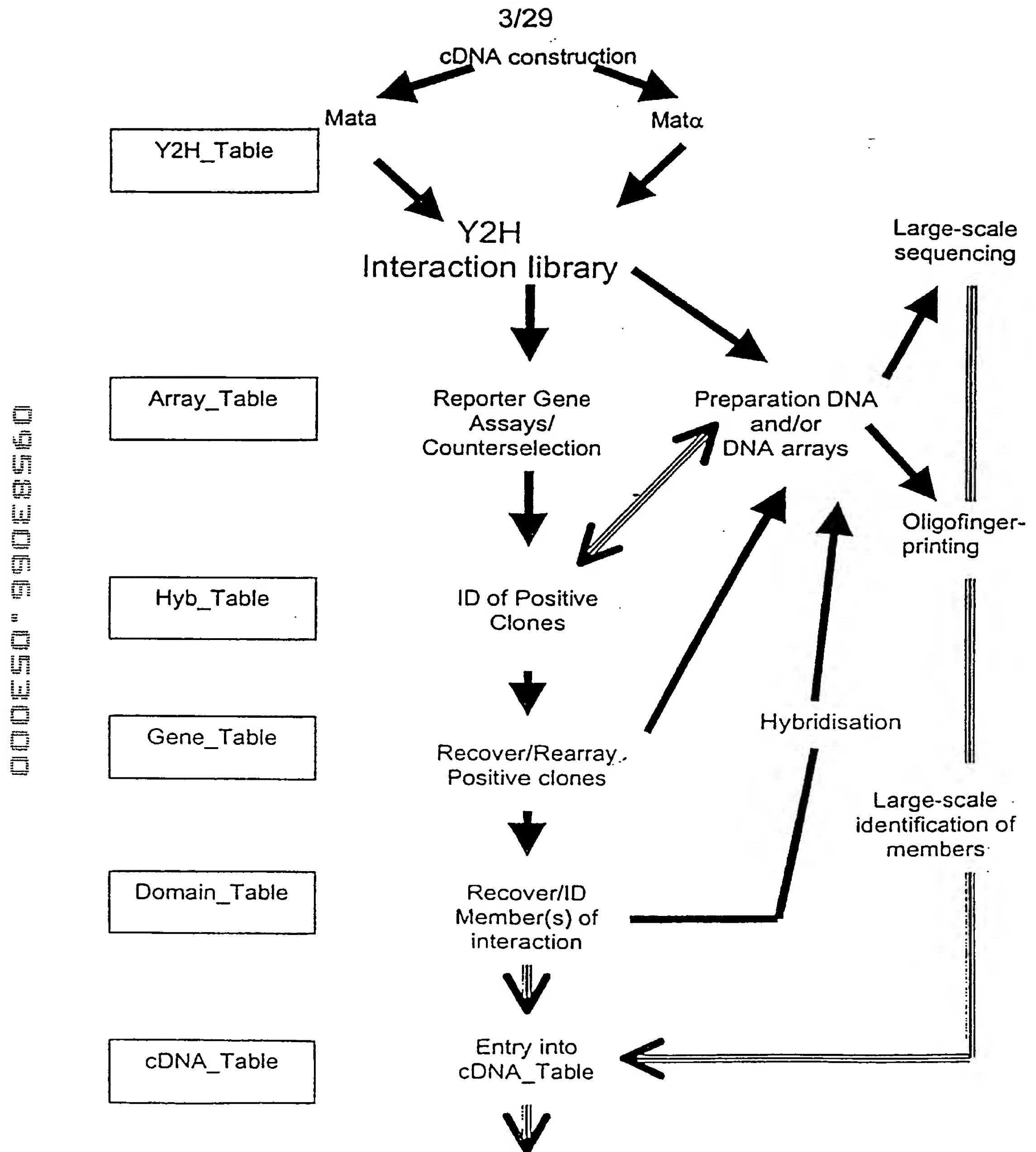


Figure 2



PathCode_Table



[illegible]

a) TCG AGT CGA CGC GGC CGC TAA CCG G

Sal I *Not* I STOP

b) TCG AGG TCG ACG CGG CCG CAG TAA CCG G

Sal I *Not* I STOP

c) TCG AGA GTC GAC GCG GCC GCT TAA CCG G

Sal I *Not* I STOP



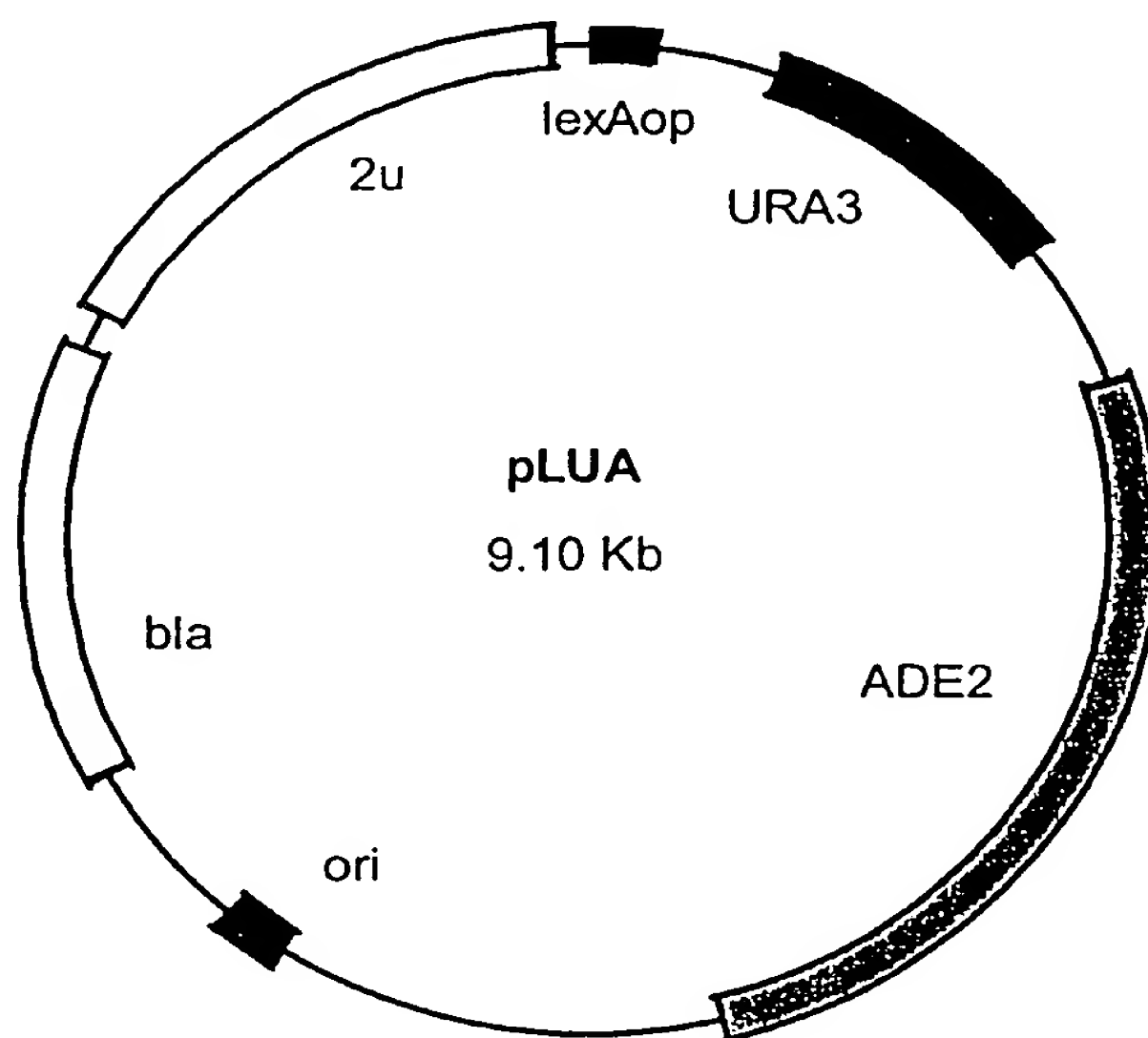
Figure 4 (II) 

Figure 5 



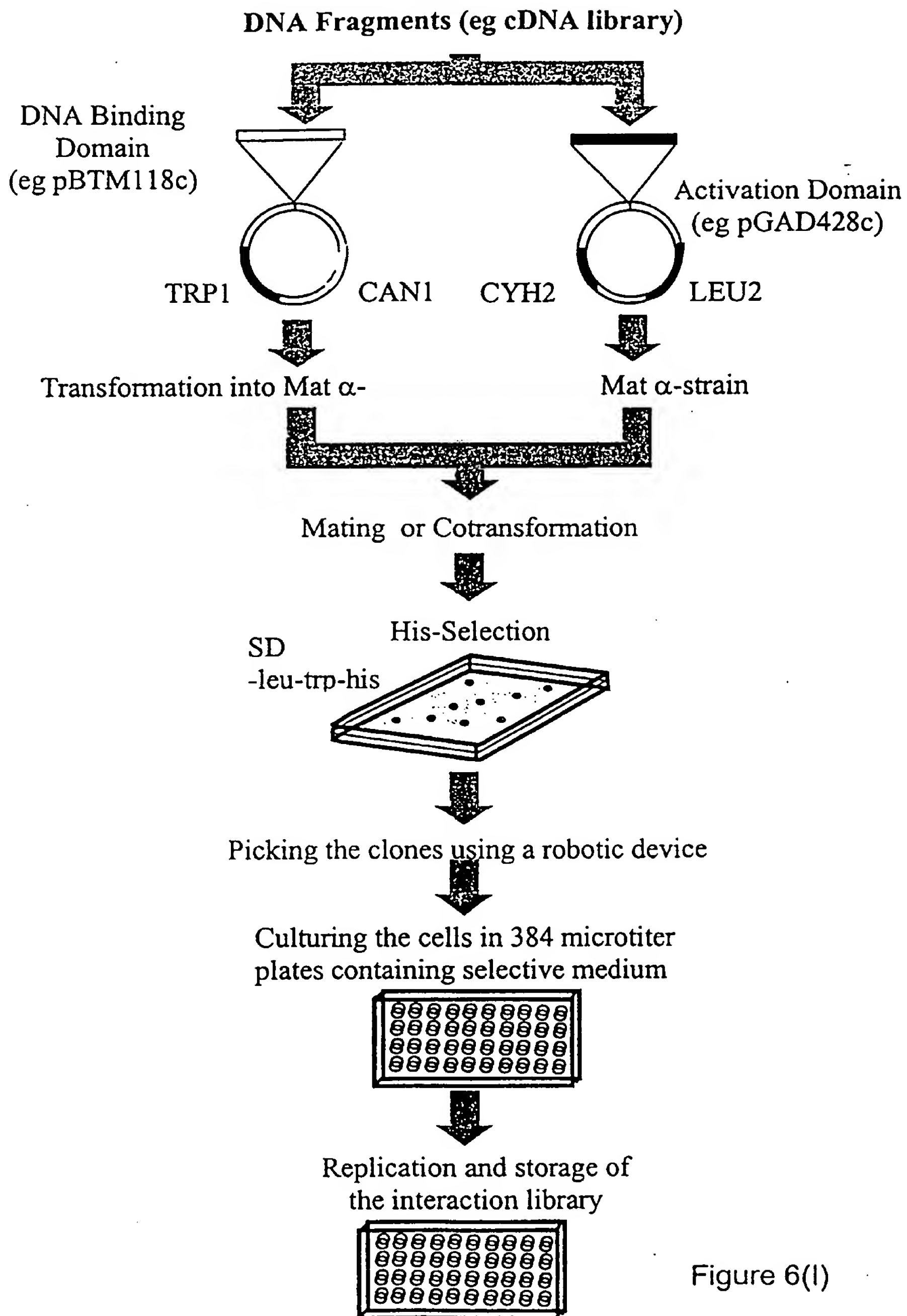
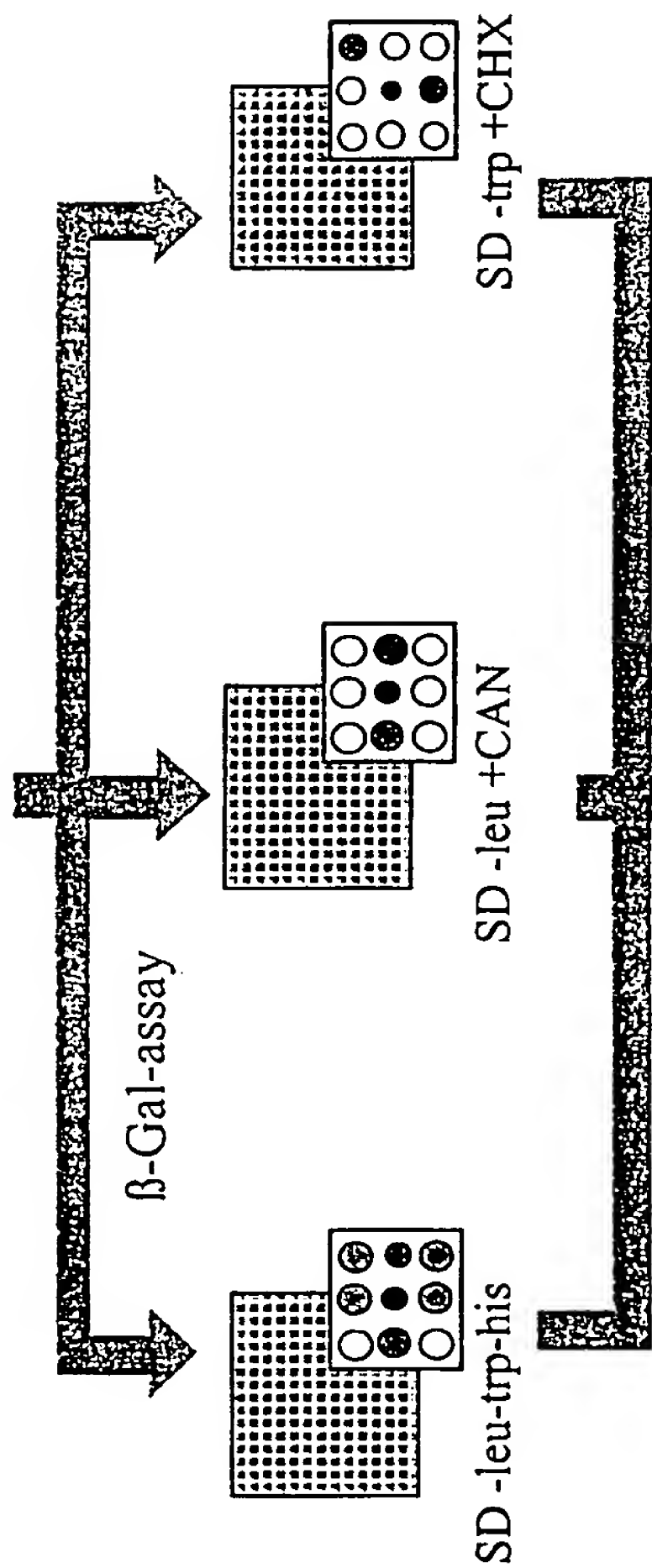


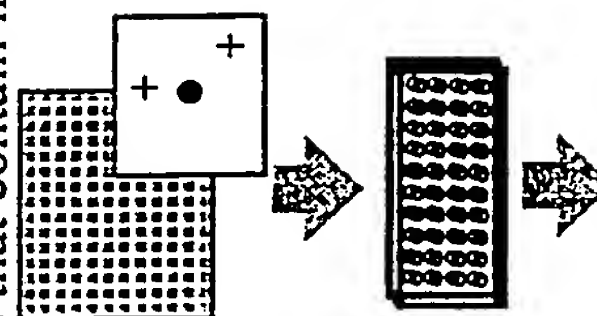
Figure 6(l)

(continuation from figure 6 (I))

Spotting of yeast cells onto a nylon membrane;
Transfer of these membranes to different selective media



Identification of clones that contain interacting molecules



(continuation see figure 6(III))

Figure 6 (II)

Identification of members comprising interactions

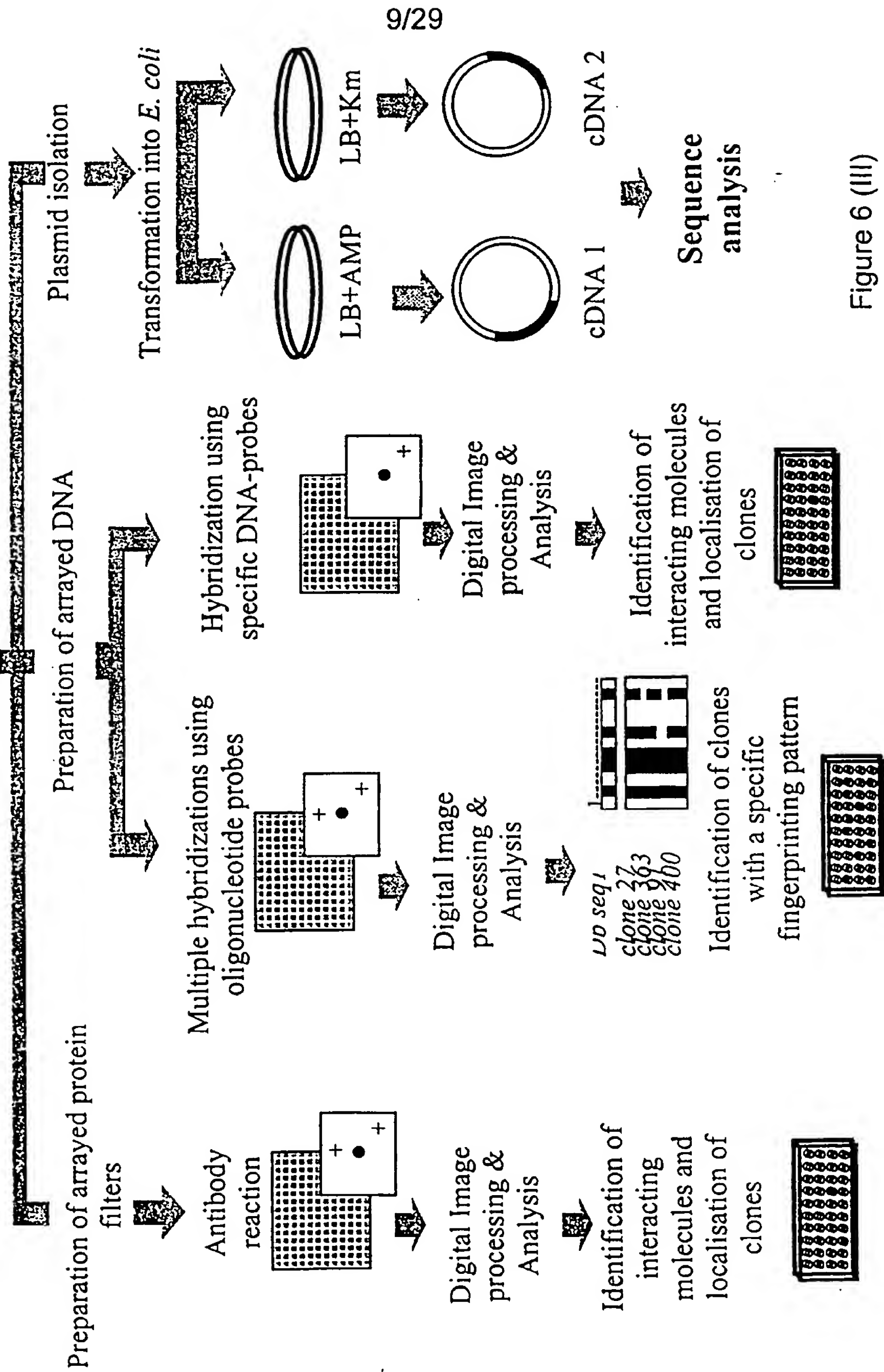
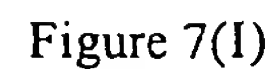
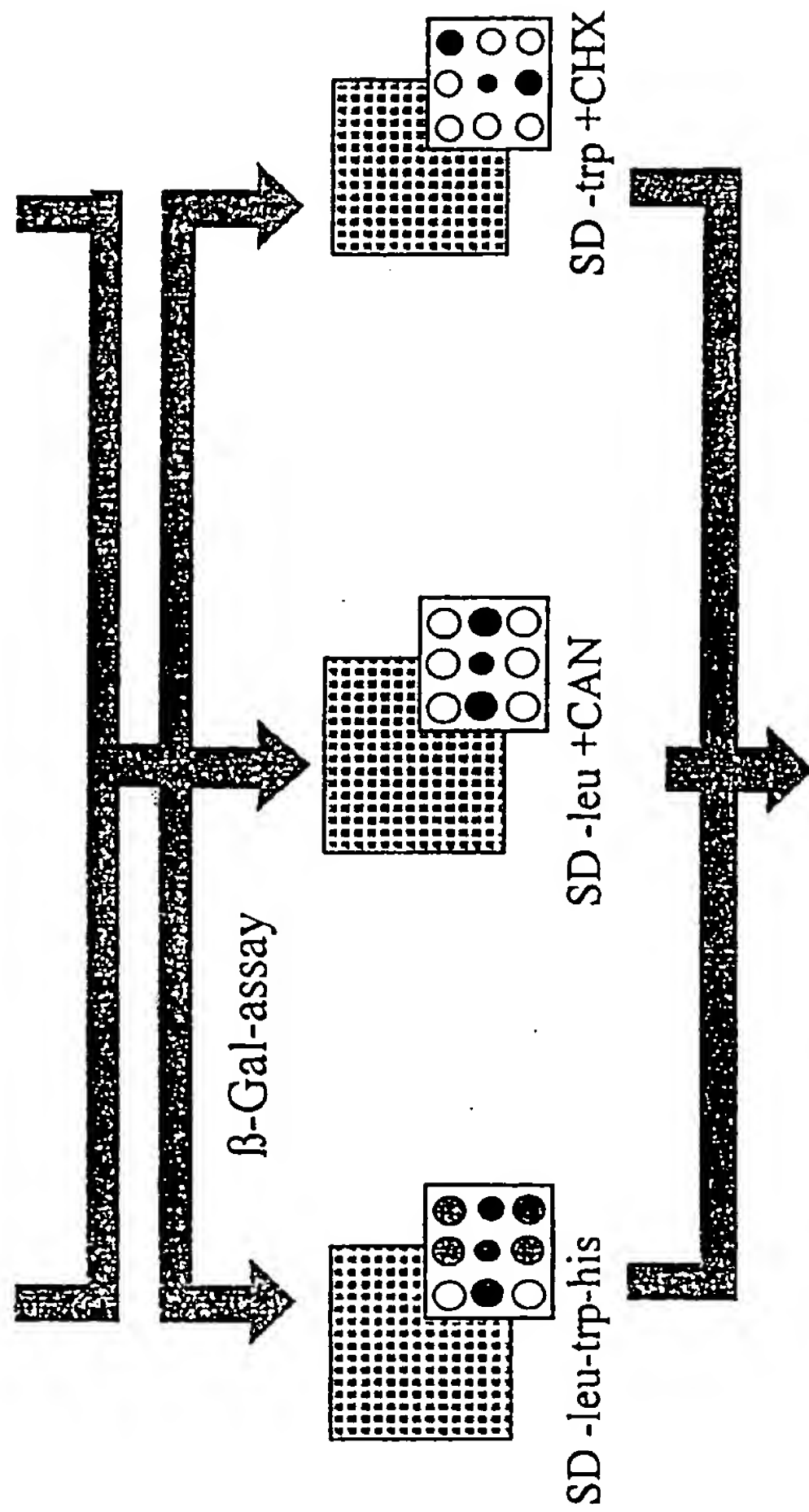


Figure 6 (III)



(continuation from figure 7 (I))



Identification of clones that contain interacting molecules

(continuation see figure 7(III))

Figure 7 (II)

Identification of genetic elements comprising interactions

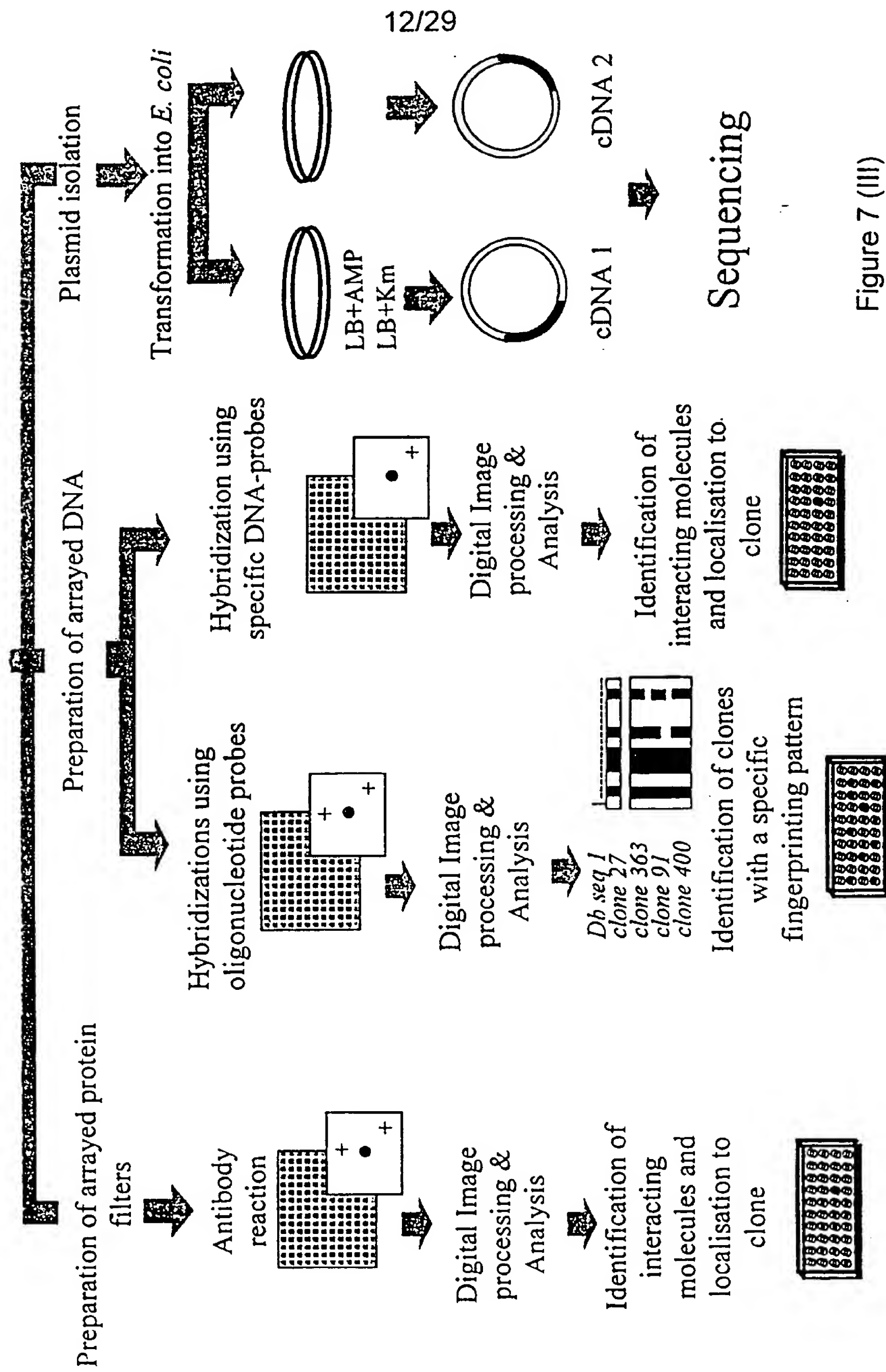


Figure 7 (III)



14/29

09583066-053000

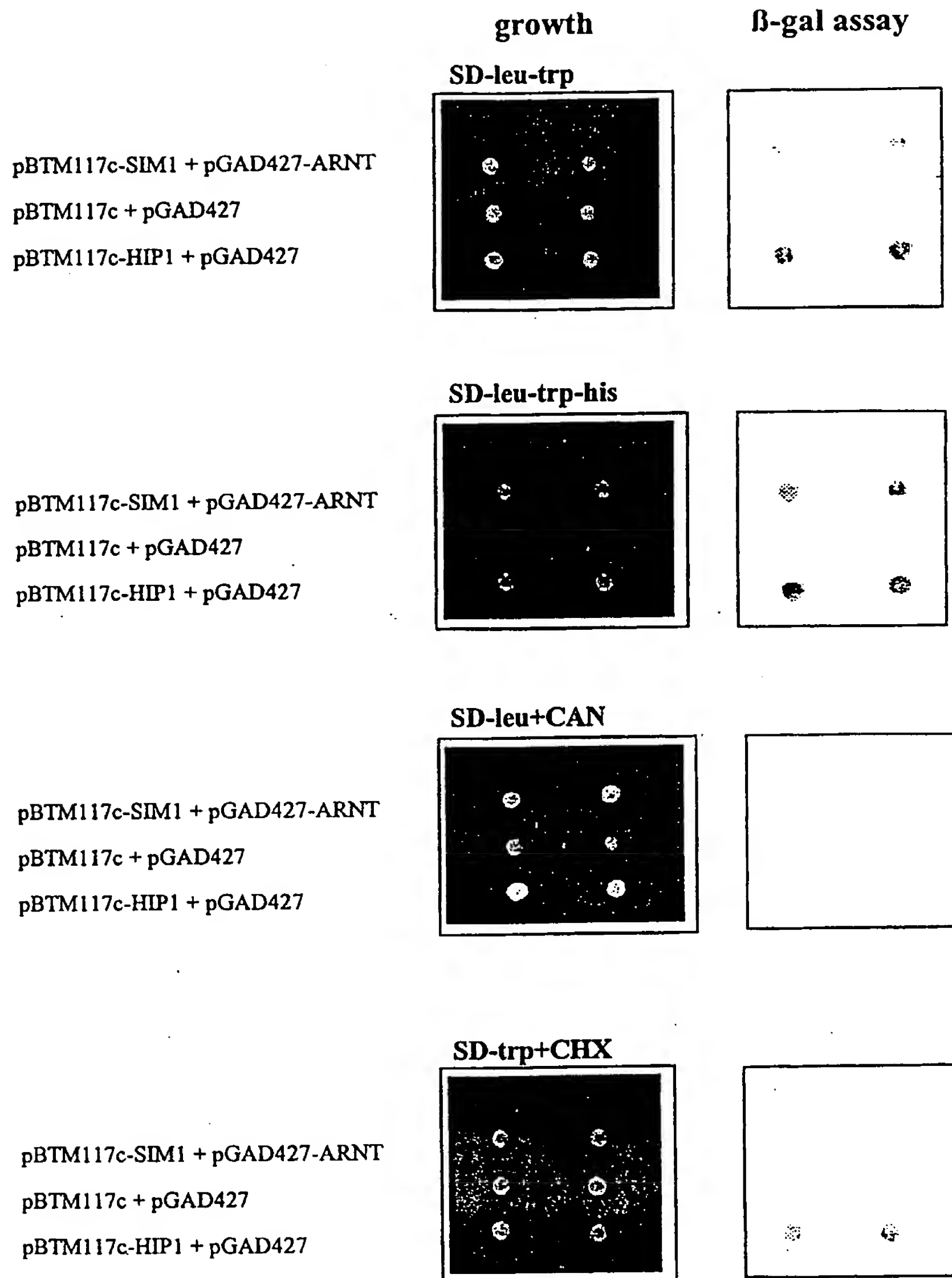


Figure 9

000E50" 990E850

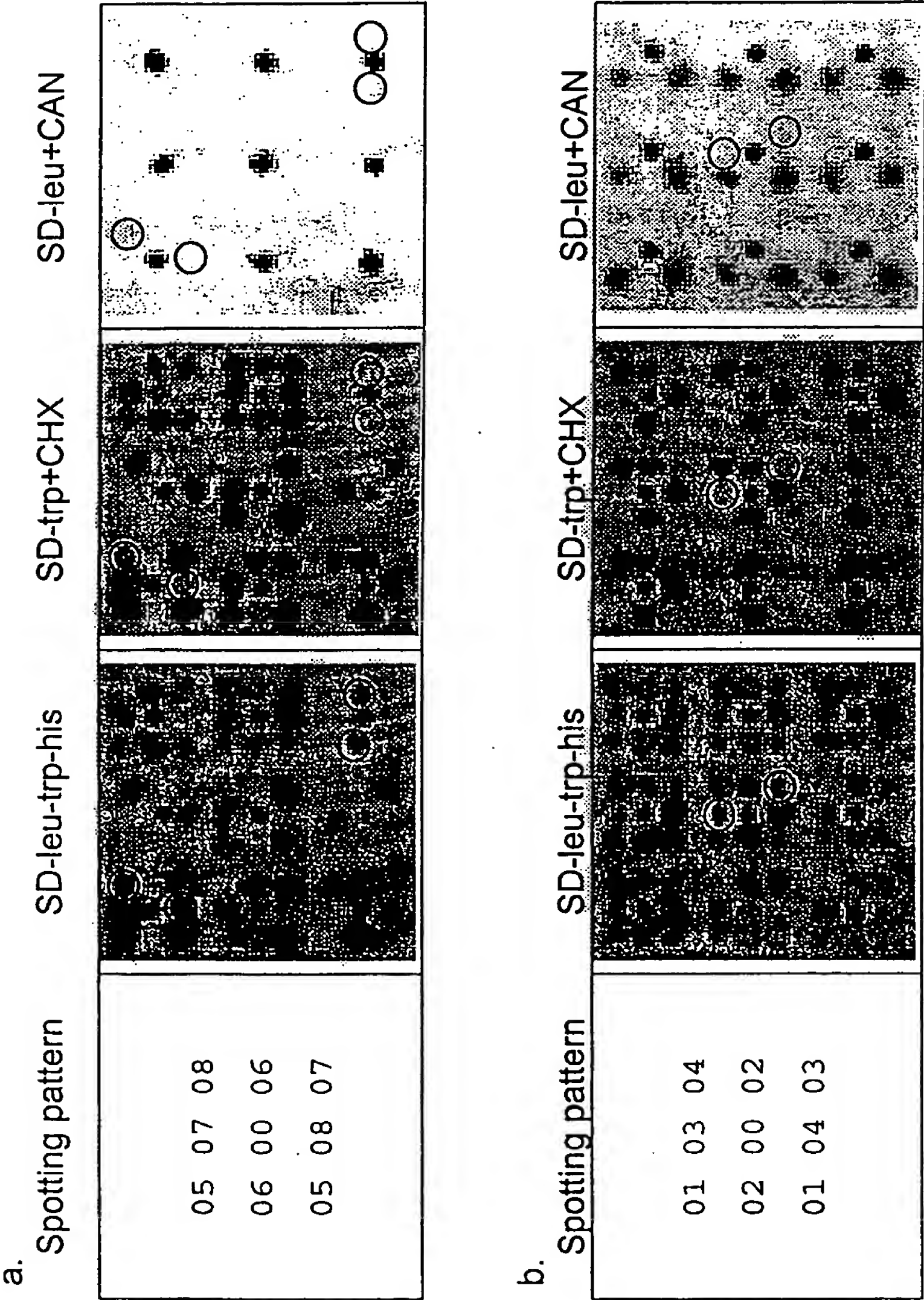


Figure 11

a.

18/29

09583065-053000

GPC - PositiveFind				
	Clone	SD-leu-trp-his	SD-trp+CHX	SD-leu+CAN
675	208	2	0	0
687	2022	2	0	0
689	2P1	3	0	0
690	2P2	2	0	0
693	2P5	3	0	0
696	2P8	3	0	0
699	2P12	3	0	0
709	2P23	2	0	0
710	2P24	2	0	0
711	3A2	3	0	0
712	3A3	3	0	0
713	3A4	3	0	0
714	3A5	3	0	0
716	3A7	3	0	0
717	3A8	3	0	0
718	3A11	3	0	0
719	3A12	2	0	0
720	3A14	3	0	0
721	3A15	3	0	0
722	3A18	2	0	0
723	3A21	2	0	0
724	3A23	2	0	0
725	3A24	3	0	0
727	3B2	3	0	0
728	3B3	3	0	0
729	3B4	3	0	0
730	3B5	3	0	0
731	3B6	3	0	0
732	3B7	3	0	0
733	3B8	3	0	0
734	3B9	3	0	0
735	3B10	3	0	0
736	3B12	3	0	0
737	3B13	3	0	0
738	3B14	3	0	0
739	3B15	3	0	0
741	3B17	3	0	0

b.

GPC - PositiveFind				
1	Clone	SD-leu-trp-his	SD-trp+CHX	SD-leu+CAN
1	Clone	SD-leu-trp-his	SD-trp+CHX	SD-leu+CAN
1733	6L22	3	0	0
2302	8N24	2	0	0

Figure 12

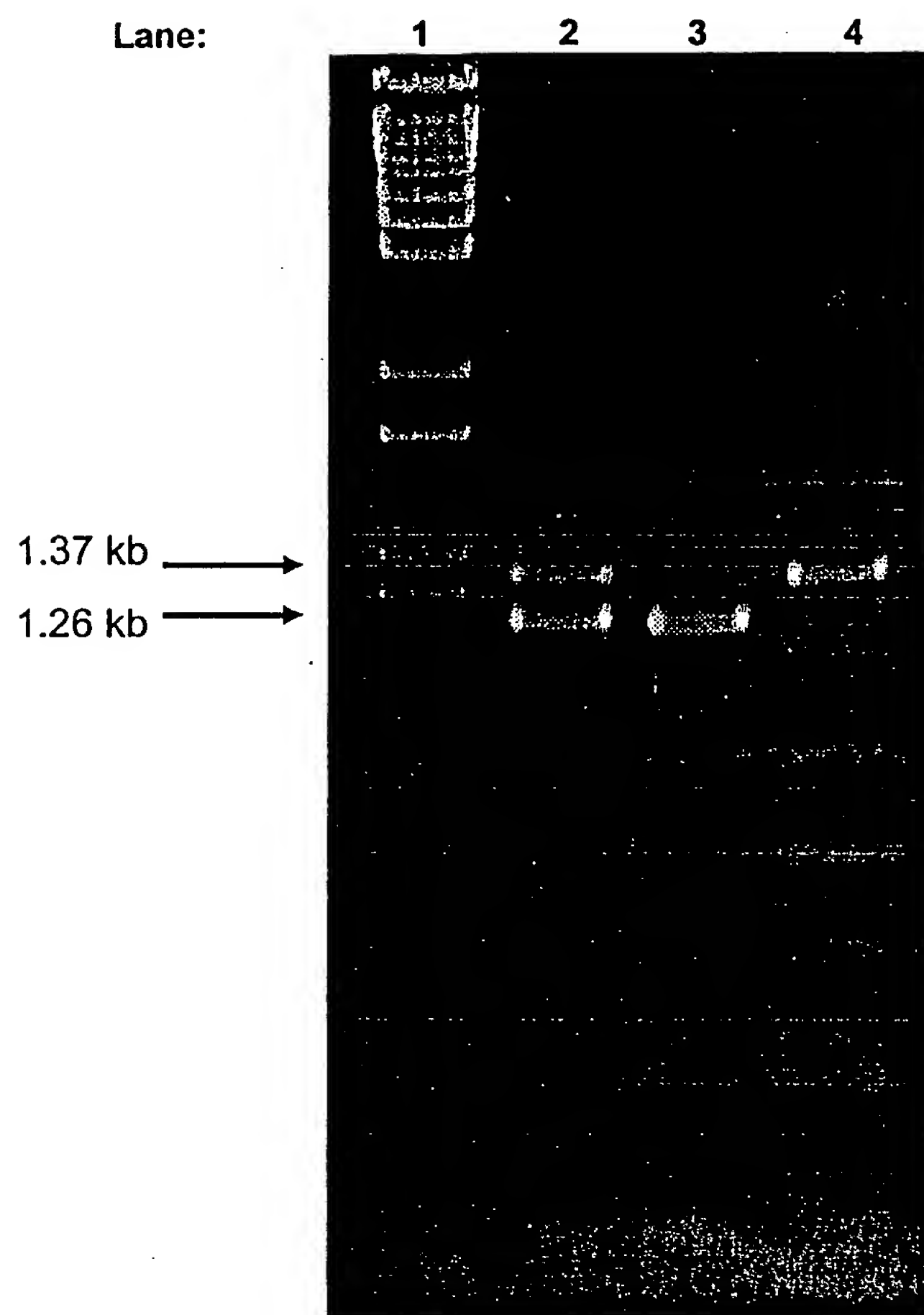


Figure 14

000050" 990E8560

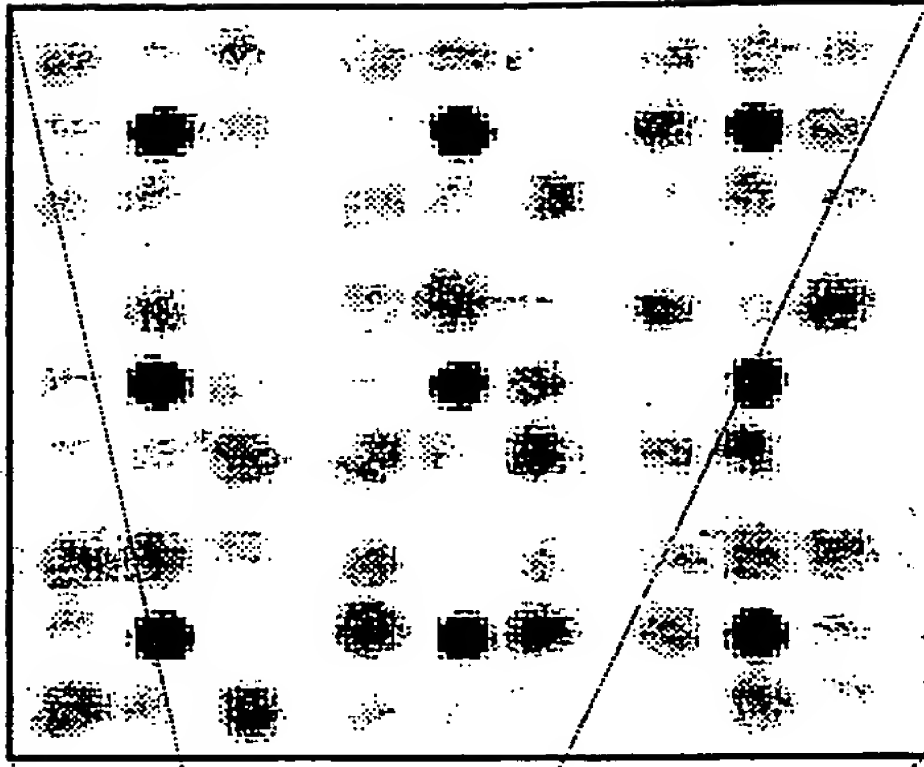
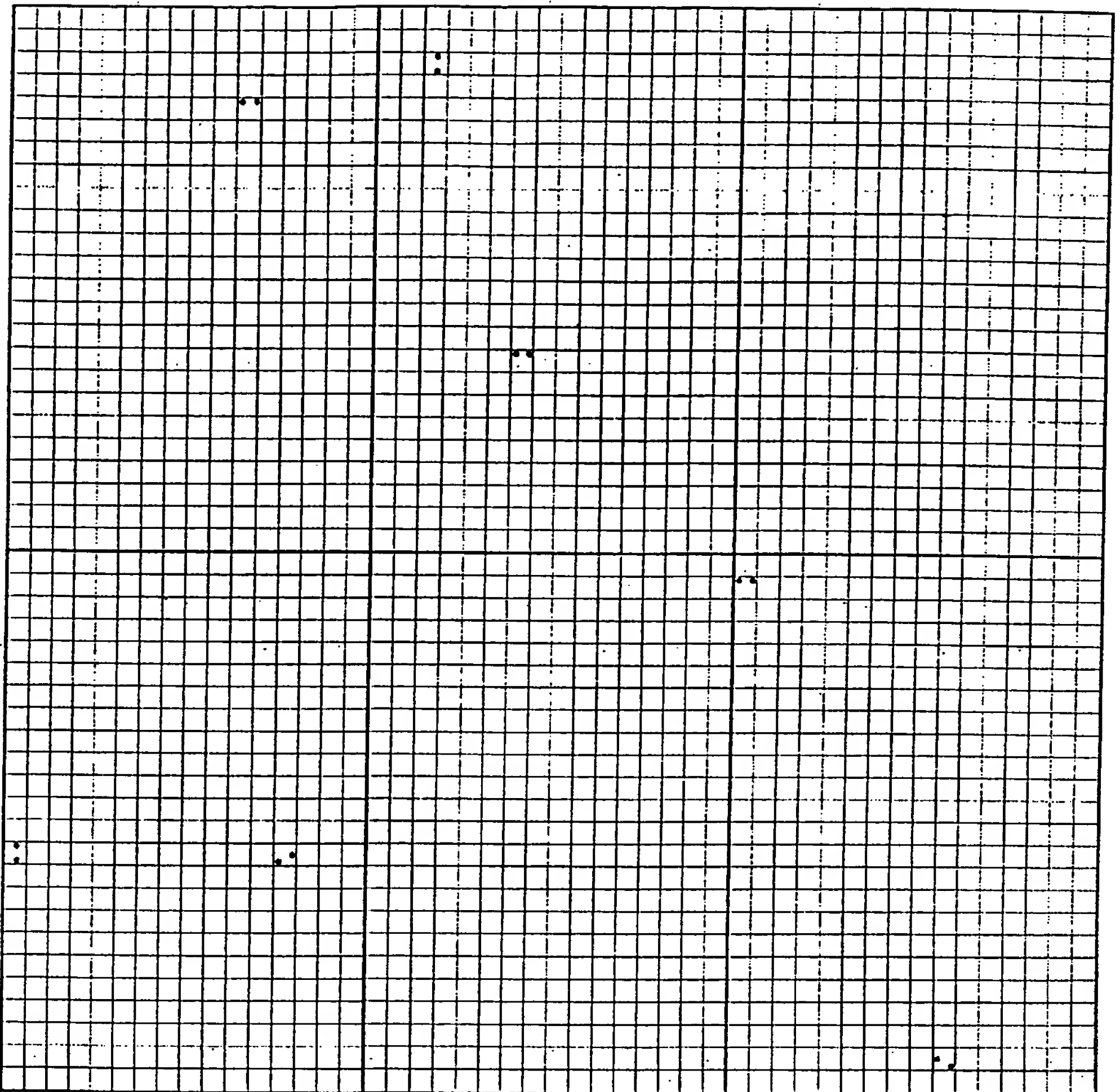


Figure 15



1st Hybridisation

Protein A: ●

Interaction Map

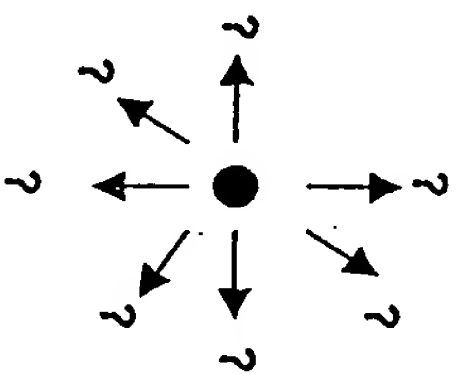
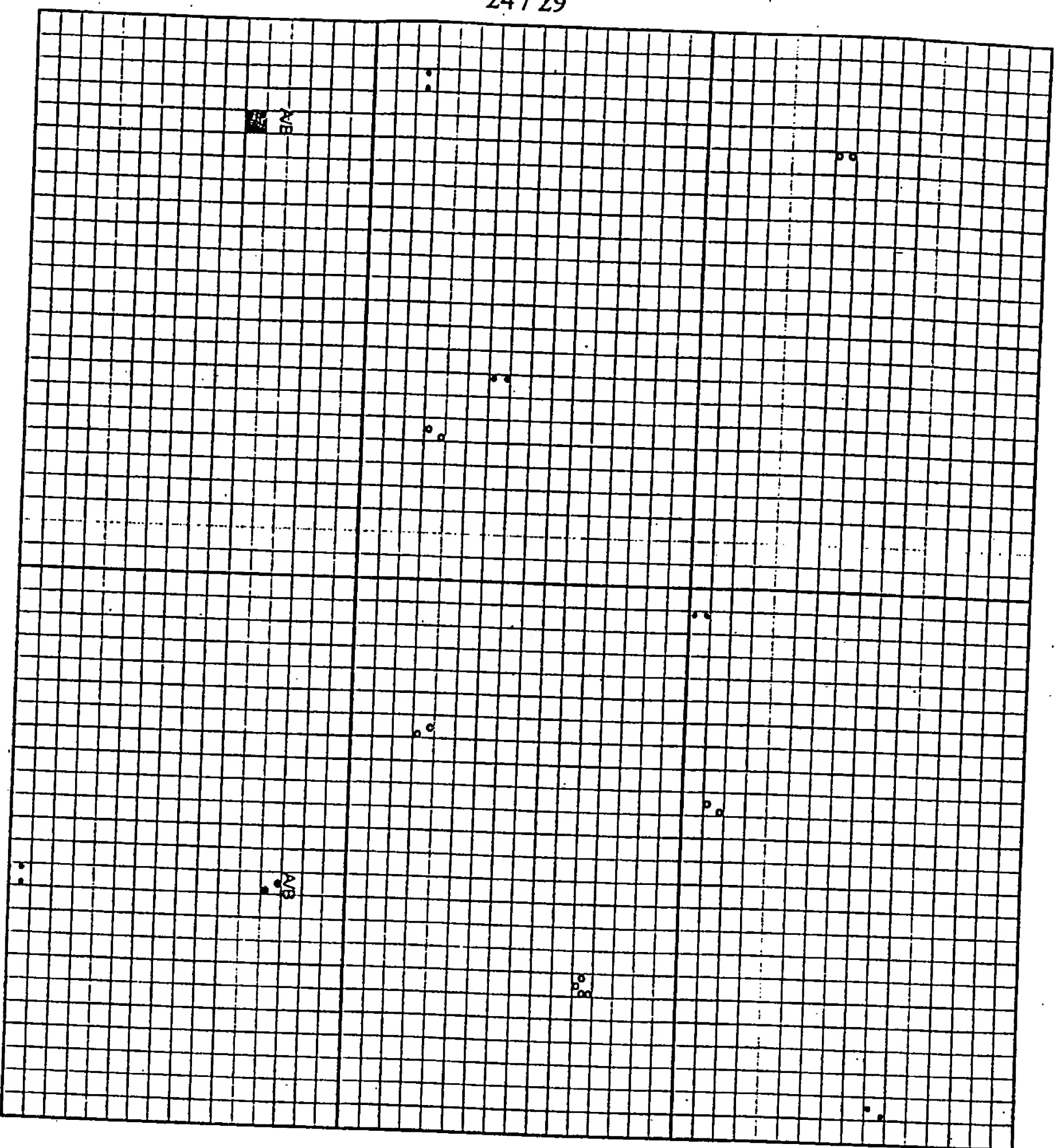


Figure 17

09583065 , 053000



1st Hybridisation

Protein A: ●

2nd Hybridisation

Protein B: ○

Interaction Map

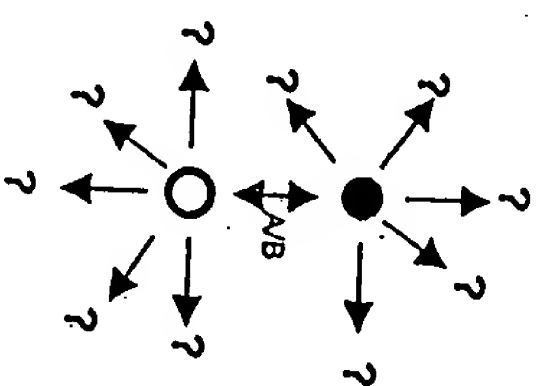


Figure 18

09583015.052001

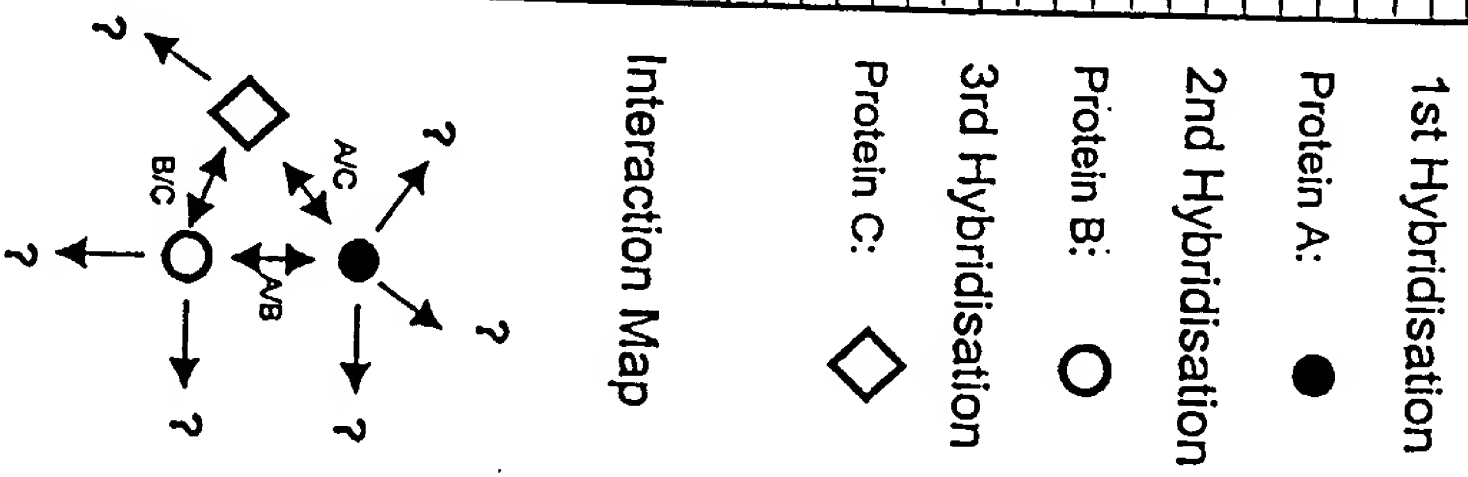
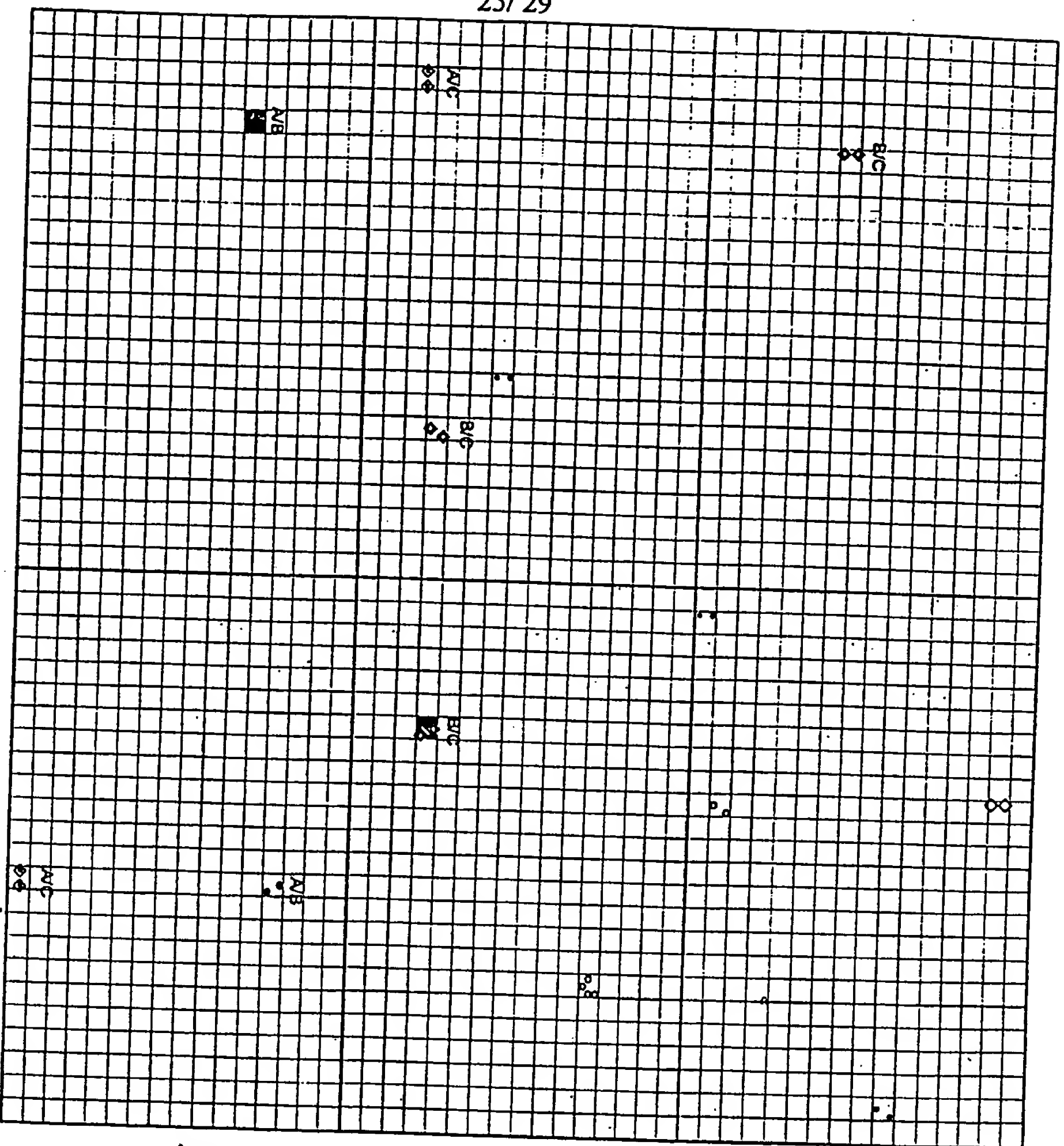


Figure 19

09583065 053000

000E50" 990E8560

26/29

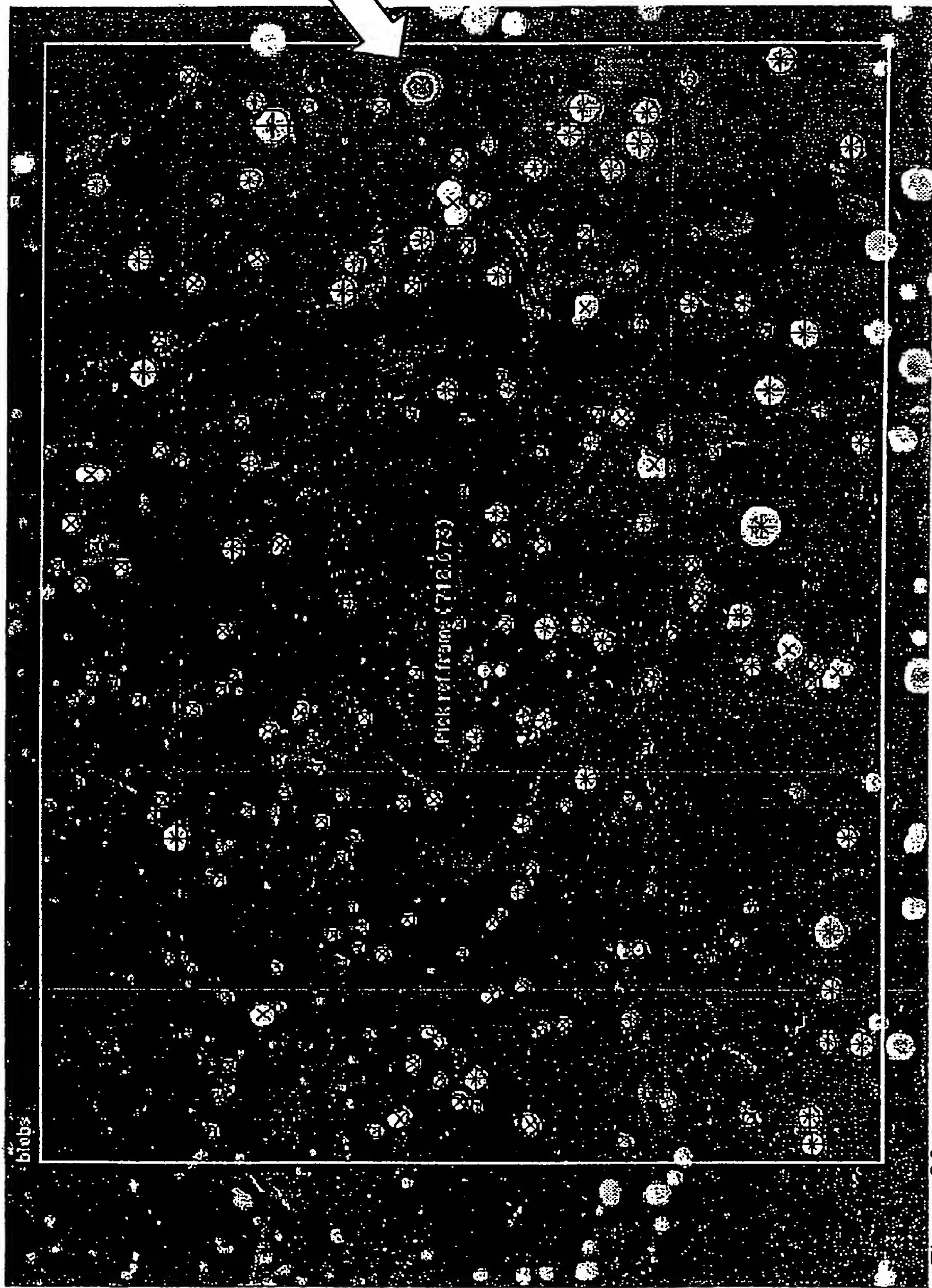


Figure 20

X-Gal



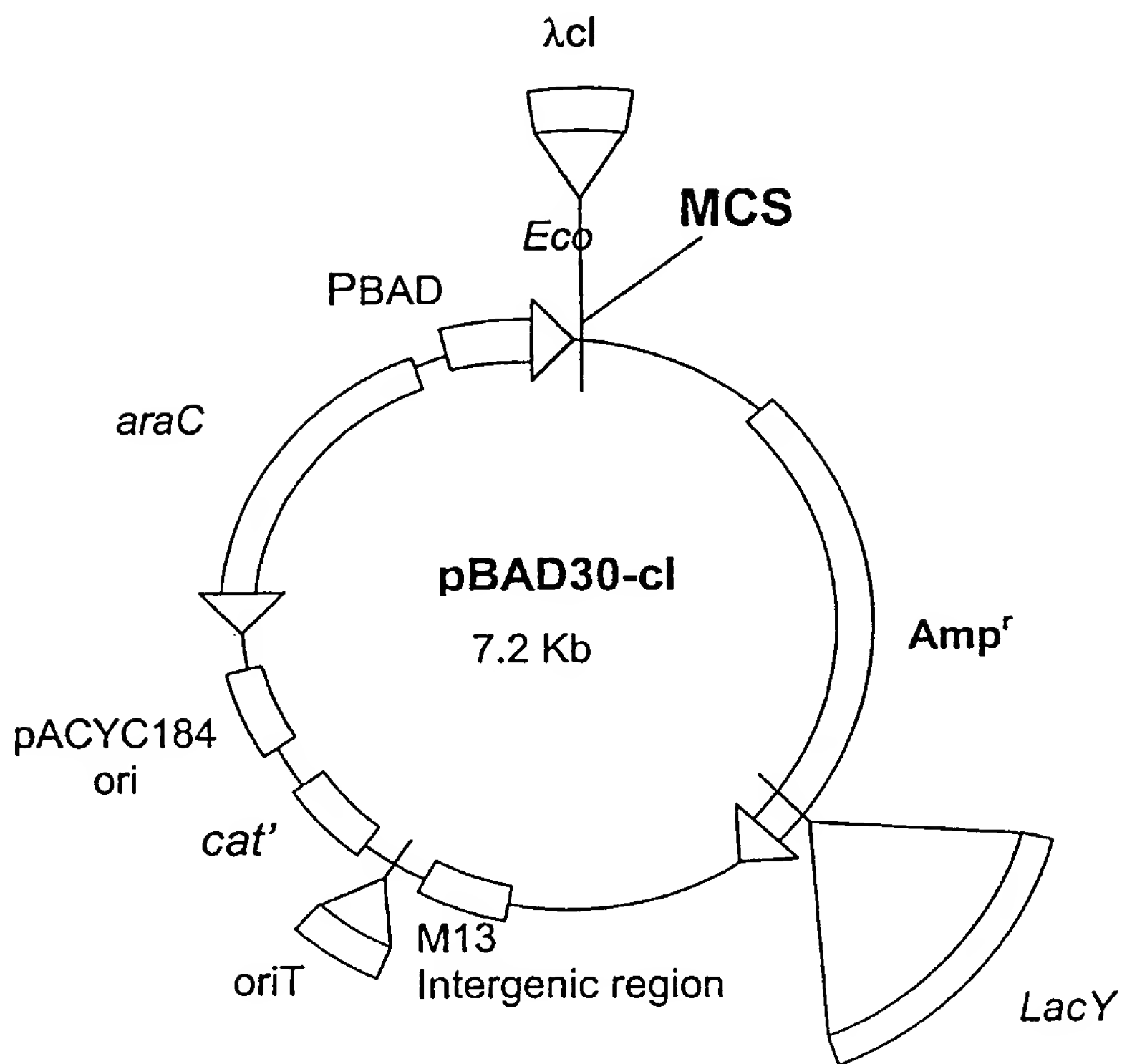


Figure 22 (I)

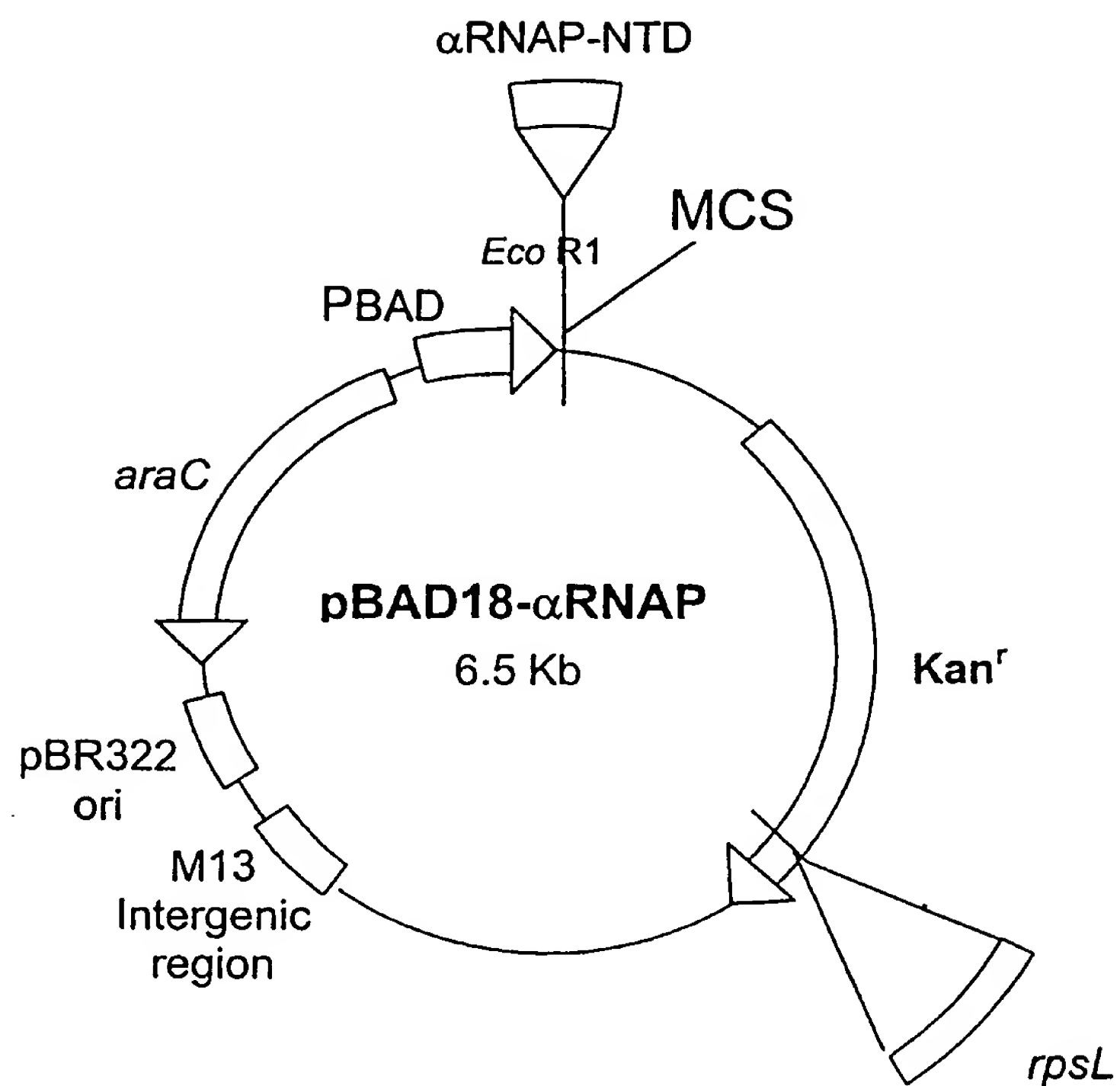


Figure 22 (II)